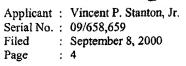
Applicant: Vincent P. Stanton, Jr. Serial No.: 09/658,659
Filed: September 8, 2000
Page: 2

	3150	2756A>G	D9190	;
	3207	2813G>T		
	3209	2815G>C	G939F	
	5444	5050C>A	3'	
	5551	5157G>A	3 '	
	5573	5179C>T	3'	
	5659	5265T>C	3'	
	5678	5284T>C	3 '	
	5874	5480C>T	3'	
	5934	5540A>G	31	
D78586	D78586	114010	GEN-BR	CAD PROTEIN
	3434	3408C>T	Silent	
	4313	4287T>C	Silent	
	4799	4773A>G	Silent	
	5255	5229C>T	Silent	
	5455	5429G>A		
	5507	5481T>C	Silent	
	5810	5784C>T	Silent	
	6128	6102C>T		
	6626	6600C>T		
	6686	6660C>T	Silent	
U09178			GEN-HA	
Dihydropyri	-	-		
	166	85T>C		
	577	496A>G		
	638	557A>G		
	1708	1627A>G		
	3432	3351T>C		
	3682	3601C>T		
	3730	3649G>A	3'	
	3925	3844A>G	3'	
	3937	3856T>C	3'	
U19720	U19720	600424 (GEN-I1	Folate
Transporter		000.7	507.	
	175	80G>A	R27H	
	341	246C>G	Silent	
	791	696C>T	Silent	
	1067 1337	972G>A	Silent Silent	
		1242C>A		, and a second second
	1997	1902T>C 2005^2006insG	3' 3'	
	2100			
	2582	2487T>G	31	
	2617	2522C>T	31	
	2652	2557T>C	3'	

Applicant: Vincent P. Stanton, Jr. Serial No.: 09/658,659
Filed: September 8, 2000
Page: 3

folate carrier (RFC1) gene, exons la, lc and lb 431 431A>G Intron 441 441A>G Intron 498 498C>T Intron 579 579G>C Intron 599 599G>C Intron 591 593 593T>C Intron 654 654 624A>C Intron 639 639A>C Intron 655 655T>C Intron 655 655T>C Intron 656 655T>C Intron 657 655T>C Intron 658 659A>C Intron 732 732T>C Intron 732 732T>C Intron 1293 1293A>C Intron 1293 1293A>C Intron 1293 1293A>C Intron 1379 1379T>C Intron 1379 1379T>C Intron 1590 1590 1590C>T Intron 1590 1590C>T Intron 1688 1688C>G Intron 1590 1590C>T Intron 1698 1688C>G Intron 1590 1590C>T Intron 1590 1590C>T Intron 1698 1688C>G Intron 1590 1590C>T Intron 1590 1590C>T Intron 1698 1688C>G Intron 1700 1888 1885O 1885						
431					-	piens reduced
441	iolate carr		gene,			
498						
S79 S79G>C Intron						
X02308 X02308 188350 GEN-KL						
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701 701A>C Intron 716 716A>G Intron 732 732T>C Intron 1293 1293A>G Intron 1322 1322C>G Intron 1379 1379T>C Intron 1590 1590C>T Intron 1688 1688C>G Intron 2401 2401A>G Intron 2429 2429G>A Intron 2488 2488C>T Intron 2594 2594G>T Intron 2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron						•
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716 716A>G Intron 732 732T>C Intron 1293 1293A>G Intron 1322 1322C>G Intron 1379 1379T>C Intron 1590 1590C>T Intron 1688 1688C>G Intron 2401 2401A>G Intron 2429 2429G>A Intron 2488 2488C>T Intron 2488 2488C>T Intron 2594 2594G>T Intron 2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron	00p1000 00.			701A>C	Intro	n
732 732T>C Intron 1293 1293A>G Intron 1322 1322C>G Intron 1379 1379T>C Intron 1590 1590C>T Intron 1688 1688C>G Intron 2401 2401A>G Intron 2429 2429G>A Intron 2488 2488C>T Intron 2594 2594G>T Intron 2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron						
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1379 1379T>C Intron 1590 1590C>T Intron 1688 1688C>G Intron 2401 2401A>G Intron 2429 2429G>A Intron 2488 2488C>T Intron 2594 2594G>T Intron 2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron						
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2401 2401A>G Intron 2429 2429G>A Intron 2488 2488C>T Intron 2594 2594G>T Intron 2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron						
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2594 2594G>T Intron 2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron		2429		2429G>A	Intro	ı
2618 2618G>A Intron 3083 3083G>A Intron 3125 3125G>A Intron		2488		2488C>T	Intro	ı
3083 3083G>A Intron 3125 3125G>A Intron		2594		2594G>T	Intro	נ
3125 3125G>A Intron		2618		2618G>A	Intro	נ
		3083		3083G>A	Intro	נ
3212 3212C>T Intron		3125		3125G>A	Intro	מ
		3212		3212C>T	Intro	ו
3619 3619T>A Intron					Intro	n
3635 3635G>A Intron		3635		3635G>A	Intro	1
		4256		4256G>A	Intro	ו
		4256		4256G>A	Intror	ו





4898	4898A>G	Intron	
5006	5006C>T	Intron	
5062	5062G>A	Intron	
5167	5167G>A	Intron	
11069	11069A>G	Intron	
11238	11238C>T	Intron	
11293	11293T>G	Intron	
11422	11422T>C	Intron	
11686	11686C>T	Intron	
12598	12598T>C	Intron	
13171	13171T>C	Intron	
13298	13298G>A	Intron	
13645	13645T>C	Intron	
13751	13751C>A	Intron	
13782	13782T>C	Intron	
13806	13806T>C	Intron	
13813	13813T>C	Intron	
14479	14479A>G	Intron	
14546	14546^insT	Intron	
14585	14585C>T	Intron	
14729	14729G>A	Intron	
14787	14787C>T	Intron	
14795	14795G>A	Intron	
15041	15041T>C	Intron	
15343	15343G>A	Intron	
15449	15449G>A	Intron	
15502	15502G>A	Intron	
15545	15545C>T	Intron	
15589	15589A>G	Intron	
15769	15769C>T	3'	
15839	15839A>G	3 '	
16148	16148G>A	3 '	
16198	16198T>G	3 '	
16202	16202G>T	Intron	
X59618 X59618	180390 G	EN-M3 R	ibonucleotide
reductase M2 polypeptide			
128	(-67)G>A	5 '	
189	(-6) T>G	5'	
524	330C>G	Silent	
1399	1205T>A	3'	
1464	1270G>A	3'	
1636	1442C>T	3 '	
1738	1544C>T	3 '	
2259	2065T>C	3'	

Applicant: Vincent P. Stanton, Jr. Serial No.: 09/658,659
Filed: September 8, 2000
Page: 5

S72487	S72487	131222	GEN-3LD) Thymi	dine	
phosphorylase,	, partial			-		
	183	19G	>A	D7N		
4	183	319C	>T	3'		
(501	437G	>C	3'		
12	299	1135G	>A	3 '		
M58602 M58	3602 1313	222 GEN	-LUB	Thymidine		
phosphorylase,	, promoter an	nd genomic	!			
1	124	124C	>T	3'		
4	139	439G	>A	3'		
10	044	1044^ins	CT	3'		
13	331	1331G	>A	3'		
19	977	1977G	>A I	Intron		
21	149	2149G	>A I	Intron		
24	167	2467A	.>G I	Intron		
	534	2634C	>G I	Intron		
	975	2975G		ntron		
	116	3116G		Intron		
	255	3255A		Intron		
	344	3344T		Intron		
)51	4051C		Intron		
	782	4782G		Intron		
	022	5022T		ntron		
	266	5266G		Intron		
	285	5285C		Intron		
	138	5438T		Intron		
	182	5482C		Intron		
	529	5629G		Intron		
	548	5648C		intron		
	731	5731G		Intron		
M98045	M98045	136510			sapiens	
folylpolygluta						
	302	732C		Silent		
	747	1677G		3 '		
	900	1830T		3'		
		510 GEN		Human	•	
folylpolygluta	mate synthet	ase (FPGS) gene,	exons 5-11,	, and	
partial cds		14040				
	124	1424C		ntron		
	549	1649G		ntron		
	554	2554A		ntron		
U24252 U24252 136510 GEN-LUF Folylpolyglutamate synthetase, promoter and exons 1-4						
	-	_				
	263	263A		ntron		
2	266	266G	ν Γ Ι	ntron		

Applicant: Vincent P. Stanton, Jr.

Serial No.: 09/658,659

Filed: September 8, 2000

Page: 6

Attorney's Docket No.: 11926-015001 / 0017.CIP3

5'

K27Q

527		527	1C>G	Intron			
1037		1037	1037A>G				
	1139		1139G>A				
	1217	1217	7C>T	Intron			
	1647	1647	7C>T	Intron			
	1955		G>A	Intron			
	2017	2017	/G>A	Intron			
	2037	2037	7G>A	Intron			
	2189	2189	A>G	Intron			
	2282	2282	C>T	Intron			
	2309		A>G	Intron			
U09806	U09806	236250	GEN-41	FZ H	uman		
	etrahydrofolat					(SEQ	ID
NO:1)	•		·	•			
,	120	120	T>C	Silent			
	464	464	T>G	M155R			
	519		519C>T				
	668	668	668C>T				
	1059		1059T>C				
	1289		1289C>A				
1308			1308T>C				
	1784		IG>A	3' 3'			
AF061655	AF061655	123920	GEN-LUJ	Cyt	idine		
deaminase,	promoter						
	575	575	T>C	Intron			
	648	648	BT>C	Intron			
771		771	LG>C	Intron			
	883	883	3G>A	Intron			

In the Claims:

Please amend claims 182-201 as follows:

941

1051

182. (Amended) An isolated nucleic acid probe comprising at least 15 contiguous

941^insC

1051A>C

nucleotides of the nucleotide sequence of SEQ ID NO:1 (methylenetetrahydrofolate reductase), the probe comprising at least one of:



